

OIPE

RAW SEQUENCE LISTING DATE: 03/04/2002 PATENT APPLICATION: US/10/077,438 TIME: 14:44:16

Input Set : A:\A080pct.txt

4 <110 - APPLICANT: MacKay, Fabienne

Browning, Jeffrey

5

Output Set: N:\CRF3\03042002\J077438.raw

```
Ambrose, Christine
              Ischopp, Jurg
              Schneider, Pascal
              Thompson, Jeffrey
     10
              Biogen, Inc.
     11
              Apotech R&D S.A.
     13 <120 - TITLE OF INVENTION: Baff Receptor (BCMA), An
              Immunoregulatory Agent
     16 <130 - FILE FEFERENCE: A080PCT
C--> 18 <140> CURRENT APPLICATION NUMBER: US/10/077,438
C--> 19 <141> CURRENT FILING DATE: 2002-02-18
     21 - 150 - PRIOR APPLICATION NUMBER: 60/149,378
     21 - 151 - PRIOR FILING DATE: 1999-08-17
     24 - 150 - PRIOR APPLICATION NUMBER: 60/181,684
     25 -: 151 - PRIOR FILING DATE: 2000-02-11
     27 -: 150 - PFIOR APPLICATION NUMBER: 60/183,536
     28 :151 - PRIOR FILING DATE: 2000-02-18
     30 < 160 > \text{NUMBER OF SEQ ID NOS: 8}
     32 <170: SOFTWARE: FastSEQ for Windows Version 4.0
     34 -: 210: SEQ ID NO: 1
     35 - 211 > LENGTH: 184
     36 -: 212 - TYPE: PRT
     37 <213> OFGANISM: homo sapien
     39 < 400 > SEQUENCE: 1
            Leu Gin Met Ala Giy Gin tys Ser Gin Ash Giu Tyr Phe Asp
     4.1. 1
                         5
                                              10
     42 Leu Leu His Ala Cys Ile Pro Cys Gln Leu Arg Cys Ser Ser Asn Ihr
     43
                     20
                                          25
                                                               30
     44 Pro Pro Leu Thr Cys Gln Arg Tyr Cys Asn Ala Ser Val Thr Asn Ser
     15
                35
                                     40
                                                          45
     46 Val Lys Gly Thr Asn Ala Ile Leu Trp Thr Cys Leu Gly Leu Ser Leu
                                 55
     48 Ile Ile Ser Leu Ala Val Phe Val Leu Met Phe Leu Leu Arg Lys Ile
                             70
                                                  75
     50 Ser Ser Glu Pro Leu Lys Asp Glu Phe Lys Asn Thr Gly Ser Gly Leu
                         85
                                              90
     52 Leu Gly Met Ala Asn Ile Asp Leu Glu Lys Ser Arg Thr Gly Asp Glu
     5 3
                    100
                                          105
     54 lle lle Leu Pro Arg Gly Leu Glu Tyr Thr <mark>Val</mark> Glu Glu Cys Thr Cys
     ۲, ۲
        115
                                     120
     56 Glu Asp Cys Ile Lys Ser Lys Pro Lys Val Asp Ser Asp His Cys Phe
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```
57 130
                           135
                                                140
58 Pro Leu Pro Ala Met Glu Glu Gly Ala Thr Ile Leu Val Thr Ihr Lys
59 145
                       150
                                            155
6) Thr Asn Asp Tyr Cys Lys Ser Leu Pro Ala Ala Leu Ser Ala Thr Glu
                   165
                                        170
  Ile Blu Lys Ser Ile Ser Ala Arq
               180
79
უნ <210 + SEQ ID NO: 2
nd <211 - LENGIH: 552
67 ≤212 - TYPE: DNA
68 <213 · ORGANISM: homo sapien
70 <400 + SEQUENCE: 2
atgttgcaga tggctgggca gtgctcccaa aatgaatatt ttgacagttt gttgcatgct
                                                                            60
72 typearacett greaactteg atgitettet aatacteete etetaacatg teaggitatt
                                                                           120
73 qtaatqcaaq tqtqaccaat tcaqtqaaaq gaacqaatqc gattctctqq acctqtttqq
                                                                           180
74 gactgagett aataatitet tiggeagiit tegigetaat gittiigeta aggaagataa
                                                                           240
75 yotolgaaco attaaaqgao gagtttaaaa acacaggato aggtotootg ggcatggota
Th acattgacct qqaaaaqaqc aqqactgqtq atqaaattat totooqagag gootogagta
                                                                           360
To caccutggaa gaatgcacct gtgaagactg catcaagagc aaaccgaagg togactctga
                                                                           420
T8 ceatingetit ecaetoccaq etaiggagga aggeqeaace atteigteac cangaaaacq
                                                                           480
                                                                           540
Ty aatgactatt gcaagageet gecagetget tigagigeta eggagataga gaaatcaatt
                                                                           552
80 totqutaggt aa
82 <210 - SEQ ID NO: 3
83 <211 + LENGTH: 207
84 -: 212 - TYPE: PRT
85 <213 · ORGANISM: homo sapien
87 <400 · SEQUENCE: 3
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                                        10
90 Thr Gly Met Leu Gln Met Ala Gly Gln Cys Ser Gln Asn Glu Tyr Phe
91
               20
                                    25
92 Asp Ser Leu Asp Val Thr Met Leu Gln Met Ala Gly Gln Cys Ser Gln
GB
                               40
                                                    45
94 Asn Glu Tyr Phe Asp Ser Leu Leu His Ala Cys Ile Pro Cys Gln Leu
G7 65
                                            75
                       70
98 Pro Cys Gln Leu Arg Cys Ser Ser Asn Thr Pro Pro Leu Thr Cys Gln
94
                                        90
100 Arg Tyr Cys Asn Ala Ser Val Thr Asn Ser Val Lys Gly Gln Arg Tyr
1(-1
                100
                                     105
                                                         110
162 Cys Asn Ala Ser Val Thr Asn Ser Val Lys Gly Val Asp Lys Thr His
           115
                                120
                                                     125
104 Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val
       130
                            135
106 Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr
                        150
                                            155
108 Pro Glu Val Ihr Cys Val Val Val Asp Val Ser His Glu Asp Pro Glu
109
                                        170
                                                             175
                    165
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DATE: 03/04/2001 FATENT APPLICATION: US/10/077,438 TIME: 14:44:16

Input Set A:\A080pct.txt Output Set N:\CRF3\03042002\J077438.raw

1	 Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn A 1 180 185 190 	la Lys							
	2 Thr Lys Fro Arg Glu Glu Glu Ijr Asn Ser Iyr Val Val Ser V	al							
	5 <2105 SEQ ID NO: 4								
	<211 · LENGTH: 549								
1:7	7 <212 - TYPE: DNA								
11년	ਰ <213 - ORGANISM: homo sapien								
120	o (400) SEQUENCE: 4								
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122	l gabytsabya tyttybagat yybtyggbag tybtobbaaa atgaatatti ty	acagtttg 120							
123	+ tigoatycti geatacetty towactioga tyticitota atactector to								
124	4 cagoyttatt gtaatgeaag titjaceaat teagtgaaag gagtegacaa aa								
	3 tgeomadegt geodag med tgaadteetg gygggadegt dagtetteet et								
	κ awardewaagg acadeeteat gatetooogg acceptgagg toacatgegt gg								
	Tigtgageeacg aagaccetga ggteaagtte aactggtaeg tigaeggegt gg								
	B aatgebaaga baaagebyer ggaggagbag tabaabagba ogtabogtgt gg								
	 otcacegtee tycaccagga etggetgaat ggcaaggagt acaagtgcaa gg 	tetecaae 540							
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	2 <211; LENGTH 14)								
	→ <212: TYPE: PRT								
	4 <213 ORGANISM: homo sapien								
	n <400: SEQUENCE: 5								
	Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr L								
138									
	4 Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr I 5 20 25 30	ie ser							
140		ro Dro							
14.	l Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu P $_2$ 35 40 45	to Pio							
14 +	\pm Ser Arg Asp Glu Leu Thr Lys Asn Gln Val Ser Leu Thr Cys Le	eu Val							
144									
	5 Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser A								
	• • • 5 70 75	80							
	7 Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp So								
148									
	" GIT Ser Phe Phe LTS LTS Leu THI val ASP LTS Ser Alg Tip 6.	III OIII							
150	0 = 100 105 110 1 Glv Ash Val Phe Ser Cys Ser Val Met His Glu Ala Leu His As	an Hia							
	•	on his							
152	2 115 120 125 3 Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys								
154	•								
	6 <210 SEQ ID NO: 6								
	/ (211) LENGIH: 369 3 (212) TYPE: DNA								
	9 <213> ORGANISM: homo sapien								
	1 :400 > SEQUENCE: 6								
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	63 ccacaggigt acaccotyce eccateeegg gatgagetga ccaagaacca ggteageetg								
	163 ccaraggigt acadecique eccatecegy gaiqagetga ccaaqaacca ggicageetg (120- 164 acetgeetgg tcaaaqqett ctaleceage gacalegeeg tggaqtggga gagcaatggg (180-								

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165	cappeduada acaactacaa dabbacqcot occytyttyd actorgacyd otgottotto	240
160	protacados ayerbaccar qgacaagago agdtggcago aqdqgaacdt offict atuc	100
167	tecytyatys atgaggetet geacaascae tacacycaga agagsetete cetytotosc	360
163	gggaaatga	369
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1"2	+212 + TYPE PRT	
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17,	Met Leu Gln Met Ala Gly Gln Cys Ser Gln Asn Glu Tyr Phe Asp Ser	
1	1 5 10 15	
178	Leu Leu His Ala Cys Ile Pro Cys Gln Leu Arg Cys Ser Ser Asn Thr	
179	20 25 30	
180	Pro Pro Leu Thr Cys Glm Arg Tyr Cys Asn Ala Ser Val Thr Asn Ser	
181	35 40 45	
18.	Val Lys Gly Thr Ash Ala Ile Leu Trp Thr Cys Leu Gly Leu Ser Leu	
183	50 55 60	
184	Ile Ile Ser Leu Ala Val Phe Val Leu Met Phe Leu Leu Arg Lys Ile	
185		
185	Ser Ser Glu Pro Leu Lys Asp Glu Phe Lys Asn Thr Gly Ser Gly Leu	
187	35	
	Leu Gly Met Ala Asn Ile Asp Leu Glu Lys Ser Arg Ihr Gly Asp Glu	
189	199 105 110	
190	lle lle Leu Pro Arg Gly Leu Glu Tyr Thr Val Glu Glu Cys Thr Cys	
191	115 120 125	
192	Glu Asp Cys Ile Lys Ser Lys Pro Lys Val Asp Ser Asp His Cys Phe	
193	130 135 140	
194	Pro Leu Pro Ala Met Glu Glu Gly Ala Thr Ile Leu Val Thr Thr Lys	
	145 150 155 160	
196	Thr Asn Asp Tyr Cys Lys Ser Leu Pro Ala Ala Leu Ser Ala Thr Glu	
197	165 170 175	
198	lle Glu Lys Ser Ile Ser Ala Arg	
199	180	
	+:210.+ SEQ ID NO: 8	
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	HI212 - TYPE: DNA	
	<pre><:213 ** ORGANISM: Homo Sapien</pre>	
	::400> SEQUENCE: 8	
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208	acadagadag deceegtaag aadecadgaa geaggegaag ticatigite teaacatiet	120
	agetgetett getgeattig etetggaatt etigtagaga taltaetigt eetteeagge	180
	tgttetttet gtageteest tgttttettt ttgtgateat gttgsagatg getgggsagt	240
	gotoccaaaa tgaatatttt gacagtttgt tgcatgcttg cataccttgt caacttcgat	300
	gthettetaa taeteeteet etaacatgte agegttattg taatgeaagt gtgaccaatt	360
213	caytgaaaqq aacgaatgcg attototgga cotgtttggg actgaqotta ataatttott	420
	tggcagtttt cgtqctaatg tttttgctaa ggaagataag ctctqaacca ttaaaggacg	480
215	aqtittaaaaa cacaqqatca qqtctcctgq qcatqqctaa cattqacctq qaaaaqaqca	540
	ggactqqtqa tqaaattatt cttccqaqaq qcctcqagta cacqqtqqaa qaatqcacct	600
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_18	ctatogajja	азполеавсе	atticttqtca	ccaqqaaaan	qaatqantat	tgcaagaucc	720
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220	titcqactcq	agcagtgcca	ctttaaaaat	cttttgtcag	aatagatqat	gtgtcagatc	840
221	tctttaggat	gactgtattt	ttcagttgcc	gatacayctt	tttgtcctct	aactgtggaa	900
222	actctttatq	ttagatatat	ttctctaggt	tactgttggg	agcttaatgg	tagaaacttc	960
223	cttggtttca	tgattaaagt	ctttttttt	cctga			995

VERIFICATION SUMMARY

DAIE: 03/04/2002

PATENT APPLICATION: US/10/077,438 TIME: 14:44:18

Input Set : A:\A080pct.txt

Output Set: N:\CRF3\03042002\J077438.raw

Lila Mil^7) C: Current Application Number differs, Replaced Current Application Number Lila Mil^7) C: Current Filing Date differs, Replaced Current Filing Date